1647

12/27/00

#4

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                     RAW SEQUENCE LISTING                           DATE: 12/27/2000
                     PATENT APPLICATION: US/09/622,439              TIME: 13:41:01

                     Input Set : A:\Y04-12-1.app
                     Output Set: N:\CRF3\12272000\I622439.raw

         4 <110> APPLICANT: Yamanouchi Pharmaceutical Co., Ltd.
         5 <120> TITLE OF INVENTION: A novel G protein coupled receptor protein
         7 <130> FILE REFERENCE: Y9905
C-->     9 <140> CURRENT APPLICATION NUMBER: US/09/622,439
C-->    10 <141> CURRENT FILING DATE: 2000-08-17
        12 <150> PRIOR APPLICATION NUMBER: JP P1998-060215
        13 <151> PRIOR FILING DATE: 1998-03-12
        15 <150> PRIOR APPLICATION NUMBER: JP P1999-026771
        16 <151> PRIOR FILING DATE: 1999-02-03
        18 <160> NUMBER OF SEQ ID NOS: 26
        20 <170> SOFTWARE: PatentIn Ver. 2.0
        22 <210> SEQ ID NO: 1
        23 <211> LENGTH: 1128
        24 <212> TYPE: DNA
        25 <213> ORGANISM: Homo sapiens
        27 <220> FEATURE:
        28 <221> NAME/KEY: CDS
        29 <222> LOCATION: (1)..(1125)
        30 <223> OTHER INFORMATION: SREB1
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        35   1               5                  10                  15
        37 gcc ctg ggc ctc aag ctg gcc acg ctc agc ctg ctg ctg tgc gtg agc      96
        38 Ala Leu Gly Leu Lys Leu Ala Thr Leu Ser Leu Leu Leu Cys Val Ser
        39              20                  25                  30
        41 ctg gcg ggc aac gtg ctg ttc gcg ctg ctg atc gtg cgg gag cgc agc     144
        42 Leu Ala Gly Asn Val Leu Phe Ala Leu Leu Ile Val Arg Glu Arg Ser
        43          35                  40                  45
        45 ctg cac cgc gcc ccg tac tac ctg ctg ctc gac ctg tgc ctg gcc gac     192
        46 Leu His Arg Ala Pro Tyr Tyr Leu Leu Leu Asp Leu Cys Leu Ala Asp
        47      50                  55                  60
        49 ggg ctg cgc gcg ctc gcc tgc ctc ccg gcc gtc atg ctg gcg gcg cgg     240
        50 Gly Leu Arg Ala Leu Ala Cys Leu Pro Ala Val Met Leu Ala Ala Arg
        51  65                  70                  75                  80
        53 cgt gcg gcg gcc gcg gcg ggg gcg ccg ccg ggc gcg ctg ggc tgc aag     288
        54 Arg Ala Ala Ala Ala Ala Gly Ala Pro Pro Gly Ala Leu Gly Cys Lys
        55                  85                  90                  95
        57 ctg ctc gcc ttc ctg gcc gcg ctc ttc tgc ttc cac gcc gcc ttc ctg     336
        58 Leu Leu Ala Phe Leu Ala Ala Leu Phe Cys Phe His Ala Ala Phe Leu
        59             100                 105                 110
        61 ctg ctg ggc gtg ggc gtc acc cgc tac ctg gcc atc gcg cac cac cgc     384
        62 Leu Leu Gly Val Gly Val Thr Arg Tyr Leu Ala Ile Ala His His Arg
        63         115                 120                 125
        65 ttc tat gca gag cgc ctg gcc ggc tgg ccg tgc gcc gcc atg ctg gtg     432
        66 Phe Tyr Ala Glu Arg Leu Ala Gly Trp Pro Cys Ala Ala Met Leu Val
        67     130                 135                 140
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RAW SEQUENCE LISTING DATE: 12/27/2000
PATENT APPLICATION: US/09/622,439 TIME: 13:41:01

Input Set : A:\Y04-12-1.app
Output Set: N:\CRF3\12272000\I622439.raw

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69 tgc gcc gcc tgg gcg ctg gcg ctg gcc gcg gcc ttc ccg ccg gtg ctg   480
70 Cys Ala Ala Trp Ala Leu Ala Leu Ala Ala Ala Phe Pro Pro Val Leu
71 145                 150                 155                 160
73 gac ggc ggt ggc gcc gac gag gac gcg ccg tgc gcc ctg gag cag cgt   528
74 Asp Gly Gly Gly Asp Asp Glu Asp Ala Pro Cys Ala Leu Glu Gln Arg
75                 165                 170                 175
77 ccc gac ggc gcc ccc ggc gcg ctg ggc ttc ctg ctg ctg ctg gcc gtg   576
78 Pro Asp Gly Ala Pro Gly Ala Leu Gly Phe Leu Leu Leu Leu Ala Val
79             180                 185                 190
81 gtg gtg ggc gcc acg cac ctc gtc tac ctc agc ctg ctc ttc ttc atc   624
82 Val Val Gly Ala Thr His Leu Val Tyr Leu Arg Leu Leu Phe Phe Ile
83         195                 200                 205
85 cac gac cgc cgc aag atg cgg ccc gcg cgc ctg gtg ccc gcc gtc aac   672
86 His Asp Arg Arg Lys Met Arg Pro Ala Arg Leu Val Pro Ala Val Ser
87     210                 215                 220
89 cac gac tgg acc ttc cac ggc ccg ggc gcc acc ggc cag gcg gcc gcc   720
90 His Asp Trp Thr Phe His Gly Pro Gly Ala Thr Gly Gln Ala Ala Ala
91 225                 230                 235                 240
93 aac tgg acg gcg ggc ttc ggc cgc ggg ccc acg ccg ccc gcg ctt gtg   768
94 Asn Trp Thr Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Ala Leu Val
95                 245                 250                 255
97 ggc atc cgg ccc gca ggg ccg ggc cgc ggc gcg cgc cgc ctc ctc gtg   816
98 Gly Ile Arg Pro Ala Gly Pro Gly Arg Gly Ala Arg Arg Leu Leu Val
99             260                 265                 270
101 ctg gaa gaa ttc aag acg gag aag agg ctg tac aag atg ttc tac gcc   864
102 Leu Glu Glu Phe Lys Thr Glu Lys Arg Leu Cys Lys Met Phe Tyr Ala
103         275                 280                 285
105 gtc acg ctg ctc ttc ctg ctc ctc tgg ggg ccc tac gtc gtg gcc agc   912
106 Val Thr Leu Leu Phe Leu Leu Leu Trp Gly Pro Tyr Val Val Ala Ser
107     290                 295                 300
109 tac ctg cgg gtc ctg cgg gcg ccc ggc gcc gtc ccc cag gcc tac ctg   960
110 Tyr Leu Arg Val Leu Val Arg Pro Gly Ala Val Pro Gln Ala Tyr Leu
111 305                 310                 315                 320
113 acg gcc ccc gtg tgg ctg acc ttc gcg cag gcc ggc atc aac ccc gtc  1008
114 Thr Ala Ser Val Trp Leu Thr Phe Ala Gln Ala Gly Ile Asn Pro Val
115                 325                 330                 335
117 gtg tgc ttc ctc ttc aac agg gag ctg agg gac tgc ttc agg gcc cag  1056
118 Val Cys Phe Leu Phe Asn Arg Glu Leu Arg Asp Cys Phe Arg Ala Gln
119             340                 345                 350
121 ttc ccc tgc tgc cag agc ccc cgg acc acc cag gcg acc cat ccc tgc  1104
122 Phe Pro Cys Cys Gln Ser Pro Arg Thr Thr Gln Ala Thr His Pro Cys
123         355                 360                 365
125 gac ctg aaa guc att ggt tta tga                                 1128
126 Asp Leu Lys Gly Ile Gly Leu
127     370                 375
130 <210> SEQ ID NO: 2
131 <211> LENGTH: 375
132 <212> TYPE: PRT
133 <213> ORGANISM: Homo sapiens
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RAW SEQUENCE LISTING DATE: 12/27/2000
PATENT APPLICATION: US/09/622,439 TIME: 13:41:01

Input Set : A:\Y04-12-1.app
Output Set: N:\CRF3\12272000\I622439.raw

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139 Ala Leu Gly Leu Lys Leu Ala Thr Leu Ser Leu Leu Leu Cys Val Ser
140               20                  25                  30
143 Leu Ala Gly Asn Val Leu Phe Ala Leu Leu Ile Val Arg Glu Arg Ser
144           35                  40                  45
145 Leu His Arg Ala Pro Tyr Tyr Leu Leu Leu Asp Leu Cys Leu Ala Asp
146       50                  55                  60
148 Gly Leu Arg Ala Leu Ala Cys Leu Pro Ala Val Met Leu Ala Ala Arg
149   65                  70                  75                  80
151 Arg Ala Ala Ala Ala Ala Gly Ala Pro Pro Gly Ala Leu Gly Cys Lys
152                   85                  90                  95
154 Leu Leu Ala Phe Leu Ala Ala Leu Phe Cys Phe His Ala Ala Phe Leu
155               100                 105                 110
157 Leu Leu Gly Val Gly Val Thr Arg Tyr Leu Ala Ile Ala His His Arg
158           115                 120                 125
160 Phe Tyr Ala Gln Arg Leu Ala Gly Trp Pro Cys Ala Ala Met Leu Val
161       130                 135                 140
163 Cys Ala Ala Trp Ala Leu Ala Leu Ala Ala Ala Phe Pro Pro Val Leu
164   145                 150                 155                 160
166 Asp Gly Gly Gly Asp Asp Glu Asp Ala Pro Cys Ala Leu Glu Glu Arg
167                   165                 170                 175
169 Pro Asp Gly Ala Pro Gly Ala Leu Gly Phe Leu Leu Leu Leu Ala Val
170               180                 185                 190
172 Val Val Gly Ala Thr His Leu Val Tyr Leu Arg Leu Leu Phe Phe Ile
173           195                 200                 205
175 His Asp Arg Arg Lys Met Arg Pro Ala Arg Leu Val Pro Ala Val Ser
176       210                 215                 220
178 His Asp Trp Thr Phe His Gly Pro Gly Ala Thr Gly Glu Ala Ala Ala
179   225                 230                 235                 240
181 Asn Trp Thr Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Ala Leu Val
182                   245                 250                 255
184 Gly Ile Arg Pro Ala Gly Pro Gly Arg Gly Ala Arg Arg Leu Leu Val
185               260                 265                 270
187 Leu Glu Glu Phe Lys Thr Glu Lys Arg Leu Cys Lys Met Phe Tyr Ala
188           275                 280                 285
190 Val Thr Leu Leu Phe Leu Leu Leu Trp Gly Pro Tyr Val Val Ala Ser
191       290                 295                 300
193 Tyr Leu Arg Val Leu Val Arg Pro Gly Ala Val Pro Glu Ala Tyr Leu
194   305                 310                 315                 320
196 Thr Ala Ser Val Trp Leu Thr Phe Ala Gln Ala Gly Ile Asn Pro Val
197                   325                 330                 335
199 Val Cys Phe Leu Phe Asn Arg Glu Leu Arg Asp Cys Phe Arg Ala Gln
200               340                 345                 350
202 Phe Pro Cys Cys Gln Ser Pro Arg Thr Thr Glu Ala Thr His Pro Cys
203           355                 360                 365
205 Asp Leu Lys Gly Ile Gly Leu
206       370                 375
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RAW SEQUENCE LISTING DATE: 12/27/2000
PATENT APPLICATION: US/09/622,439 TIME: 13:41:01

Input Set : A:\Y04-12-1.app
Output Set: N:\CRF3\12272000\I622439.raw

209 <210> SEQ ID NO: 3
210 <211> LENGTH: 1113
211 <212> TYPE: DNA
212 <213> ORGANISM: Homo sapiens
214 <220> FEATURE:
215 <221> NAME/KEY: CDS
216 <222> LOCATION: (1)..(1110)
217 <223> OTHER INFORMATION: SREB2
219 <400> SEQUENCE: 3

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222   1               5                  10                  15
221 ccc cta aca gcc ttt ctg aaa ctg act tcc ttg ggt ttc ata ata gga   96
225 Pro Leu Thr Ala Phe Leu Lys Leu Thr Ser Leu Gly Phe Ile Ile Gly
226             20                  25                  30
228 atc agc gtg gtg ggc aac ctc ctg atc tcc att ttg cta gtg aaa gat  144
229 Val Ser Val Val Gly Asn Leu Leu Ile Ser Ile Leu Leu Val Lys Asp
230         35                  40                  45
232 aag acc ttg cat aga gca cct tac tac ttc ctg ttg gat ctt tgc ttt  192
233 Lys Thr Leu His Arg Ala Pro Tyr Tyr Phe Leu Leu Asp Leu Cys Cys
234     50                  55                  60
236 tca gat atc ctc agg tct gca att tgt ttc cca ttt gtg ttc aac tct  240
237 Ser Asp Ile Leu Arg Ser Ala Ile Cys Phe Pro Phe Val Phe Asn Ser
238 65                  70                  75                  80
240 gtc aaa aat gac tct acc tgg act tat gga act ctg act tgc aaa gtg  288
241 Val Lys Asn Gly Ser Thr Trp Thr Tyr Gly Thr Leu Thr Cys Lys Val
242                 85                  90                  95
244 att gcc ttt ctg ggg gtt ttg tcc tgt ttc cac act gct ttc atg ctc  336
245 Ile Ala Phe Leu Gly Val Leu Ser Cys Phe His Thr Ala Phe Met Leu
246             100                 105                 110
248 ttc tgc atc agt gtc ctc acc aga tac tta gct atc gcc cat cac cgc ttc  384
249 Phe Cys Ile Ser Val Thr Arg Tyr Leu Ala Ile Ala His His Arg Phe
250         115                 120                 125
252 tat aca aag agg ctg acc ttt tgg acg tgt ctg cct gtg ctc atg  432
253 Tyr Thr Lys Arg Leu Thr Phe Trp Thr Cys Leu Ala Val Ile Cys Met
254     130                 135                 140
256 gtg tgg act ctg tct gtg gcc atg gca ttt ccc ccg gtt tta gac gtg  480
257 Val Trp Thr Leu Ser Val Ala Met Ala Phe Pro Pro Val Leu Asp Val
258 145                 150                 155                 160
260 ggc act tac tca ttc att agg gag gaa gat caa tgc acc ttc cga cac  528
261 Gly Thr Tyr Ser Phe Ile Arg Glu Glu Asp Gln Cys Thr Phe Gln His
262                 165                 170                 175
264 cgc tcc ttc agg gct aat gat tcc tta ggg ttt atg ctg ctt ctt gct  576
265 Arg Ser Phe Arg Ala Asn Asp Ser Leu Gly Phe Met Leu Leu Leu Ala
266             180                 185                 190
268 ctc atc ctc cta gcc aca cag ctt gtc tac ctc aag ctg ata ttt ttc  624
269 Leu Ile Leu Leu Ala Thr Gln Leu Val Tyr Leu Lys Leu Ile Phe Phe
270         195                 200                 205
272 gtc cac gat cgg aga aaa atg aag cca gtc cag ttt gta gca gca gtc  672
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RAW SEQUENCE LISTING DATE: 12/27/2000
PATENT APPLICATION: US/09/622,439 TIME: 13:41:01

Input Set : A:\Y04-12-1.app
Output Set: N:\CRF3\12272000\I622439.raw

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273 Val His Asp Arg Arg Lys Met Lys Pro Val Gln Phe Val Ala Ala Val
274      210                 215                 220
276 agc cag aac tgg act ttt gct ggt cct gga gcc agt gtc cag gca gct   720
277 Ser Gln Asn Trp Thr Phe Ala Gly Pro Gly Ala Ser Gly Gln Ala Ala
278 225                 230                 235                 240
280 gcc aat tgg ctg gca ggc ttt ggc agg ggt ccg gca ccc ccc acc ctg   768
281 Ala Asn Trp Leu Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Phe Leu
282                 245                 250                 255
284 ctg ggc atc agg caa aat gca aac acc aca ggc aga aga agg cta ttg   816
285 Leu Gly Ile Arg Gln Asn Ala Asn Thr Thr Gly Arg Arg Arg Leu Leu
286             260                 265                 270
288 gtc tta gac gag ttc aaa atg cag aaa aga atc agc aga atg ttc tat   864
289 Val Leu Asp Glu Phe Lys Met Gln Lys Arg Ile Ser Arg Met Phe Tyr
290         275                 280                 285
292 ata atg act ttt ctg ttt gta acc ttg tgg ggc ccc tac ctg gtg gcc   912
293 Ile Met Thr Phe Leu Phe Val Thr Leu Trp Gly Pro Tyr Leu Val Ala
294     290                 295                 300
296 tgt tat tgg aga gtt ttt gca aga ggg cct gta gta cca ggg ggt ttt   960
297 Cys Tyr Trp Arg Val Phe Ala Arg Gly Pro Val Val Pro Gly Gly Phe
298 305                 310                 315                 320
300 cta aca gct gct gtc tgg atg agt ttt gcc caa gca gga atc aat cct  1008
301 Leu Thr Ala Ala Val Trp Met Ser Phe Ala Gln Ala Gly Ile Asn Pro
302                 325                 330                 335
304 ttt gtc tgc att ttc tca aac agg gag ctg agg cgc tgt tcc agc aca  1056
305 Phe Val Cys Ile Phe Ser Asn Arg Glu Leu Arg Arg Cys Phe Ser Thr
306             340                 345                 350
308 acc ctt ctt tac tgc aga aat tcc agg tta cca agg gaa cct tac tgt  1104
309 Thr Leu Leu Tyr Cys Arg Lys Ser Arg Leu Pro Arg Glu Pro Tyr Cys
310         355                 360                 365
312 gtt ata tga                                                    1113
313 Val Ile
314     370
317 <210> SEQ ID NO: 4
318 <211> LENGTH: 370
319 <212> TYPE: PRT
320 <213> ORGANISM: Homo sapiens
322 <400> SEQUENCE: 4
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326 Pro Leu Thr Ala Phe Leu Lys Leu Thr Ser Leu Gly Phe Ile Ile Gly
327             20                  25                  30
329 Val Ser Val Val Gly Asn Leu Leu Ile Ser Ile Leu Leu Val Lys Asp
330         35                  40                  45
332 Lys Thr Leu His Arg Ala Pro Tyr Tyr Phe Leu Leu Asp Leu Cys Cys
333     50                  55                  60
335 Ser Asp Ile Leu Arg Ser Ala Ile Cys Phe Pro Phe Val Phe Asn Ser
336 65                  70                  75                  80
338 Val Lys Asn Gly Ser Thr Trp Thr Tyr Gly Thr Leu Thr Cys Lys Val
339                 85                  90                  95
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VERIFICATION SUMMARY DATE: 12/27/2000
PATENT APPLICATION: US/09/622,439 TIME: 13:41:02

Input Set : A:\Y04-12-1.app
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L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date